

=> d his

(FILE 'HOME' ENTERED AT 15:30:42 ON 09 DEC 2008)

FILE 'REGISTRY' ENTERED AT 15:30:58 ON 09 DEC 2008

E TRICARBONYL

L1 72290 S E3

FILE 'CAPLUS' ENTERED AT 15:31:12 ON 09 DEC 2008

L2 30421 S (L1 OR TRICARBONYL?)

L3 26829 S L2 AND (PY<=2003)

L4 66 S L3 AND (CANCER? OR TUMOR? OR TUMOUR?)

L5 66 DUP REM L4 (0 DUPLICATES REMOVED)

=> d stat que

L1 72290 SEA FILE=REGISTRY ABB=ON PLU=ON TRICARBONYL/BI

L2 30421 SEA FILE=CAPLUS ABB=ON PLU=ON (L1 OR TRICARBONYL?)

L3 26829 SEA FILE=CAPLUS ABB=ON PLU=ON L2 AND (PY<=2003)

L4 66 SEA FILE=CAPLUS ABB=ON PLU=ON L3 AND (CANCER? OR TUMOR? OR
TUMOUR?)

L5 66 DUP REM L4 (0 DUPLICATES REMOVED)

=> log y

=> d his

(FILE 'HOME' ENTERED AT 22:07:07 ON 09 DEC 2008)

FILE 'REGISTRY' ENTERED AT 22:07:40 ON 09 DEC 2008

L1 SCREEN 1840 AND 1918 AND 2006 AND 1996
 L2 STRUCTURE UPLOADED
 L3 QUE L2 AND L1
 L4 0 S L3
 L5 2 S L3 FULL
 L6 ANALYZE L5 1- LC : 4 TERMS

FILE 'CAPLUS' ENTERED AT 22:09:15 ON 09 DEC 2008

L7 4 S L5
 L8 4 DUP REM L7 (0 DUPLICATES REMOVED)

FILE 'REGISTRY' ENTERED AT 22:09:43 ON 09 DEC 2008

L9 SCREEN 1838 AND 1918 AND 2008 AND 1993
 L10 STRUCTURE UPLOADED
 L11 QUE L10 AND L9
 L12 0 S L11
 L13 41 S L11 FULL

FILE 'CAPLUS' ENTERED AT 22:10:29 ON 09 DEC 2008

L14 13 S L13
 L15 13 DUP REM L14 (0 DUPLICATES REMOVED)

FILE 'REGISTRY' ENTERED AT 22:11:10 ON 09 DEC 2008

L16 SCREEN 1839 AND 1918 AND 1929 AND 2006 AND 1995
 L17 STRUCTURE UPLOADED
 L18 QUE L17 AND L16
 L19 0 S L18
 L20 11 S L18 FULL

FILE 'CAPLUS' ENTERED AT 22:12:15 ON 09 DEC 2008

L21 6 S L20
 L22 6 DUP REM L21 (0 DUPLICATES REMOVED)

=> d stat que

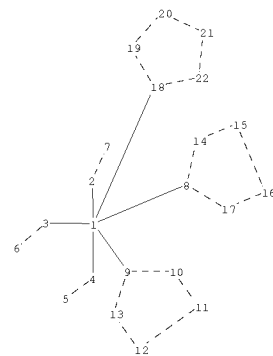
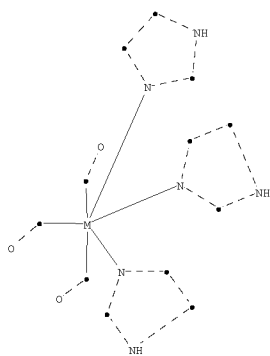
L16 SCR 1839 AND 1918 AND 1929 AND 2006 AND 1995
 L17 STR

*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***

Structure attributes must be viewed using STN Express query preparation.

L20 11 SEA FILE=REGISTRY SSS FUL L17 AND L16
 L21 6 SEA FILE=CAPLUS ABB=ON PLU=ON L20
 L22 6 DUP REM L21 (0 DUPLICATES REMOVED)

=> log y



chain nodes :

1 2 3 4 5 6 7

ring nodes :

8 9 10 11 12 13 14 15 16 17 18 19 20 21 22

chain bonds :

1-2 1-3 1-4 1-8 1-9 1-18 2-7 3-6 4-5

ring bonds :

8-14 8-17 9-10 9-13 10-11 11-12 12-13 14-15 15-16 16-17 18-19 18-22 19-20 20-21 21-22

exact/norm bonds :

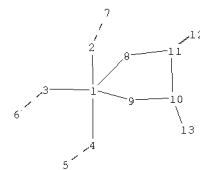
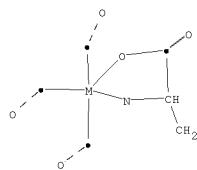
2-7 3-6 4-5 8-14 8-17 9-10 9-13 10-11 11-12 12-13 14-15 15-16 16-17 18-19 18-22 19-20 20-21 21-22

exact bonds :

1-2 1-3 1-4 1-8 1-9 1-18

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom
12:Atom 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom
22:Atom



chain nodes :

2 3 4 5 6 7 12 13

ring nodes :

1 8 9 10 11

chain bonds :

1-2 1-3 1-4 2-7 3-6 4-5 10-13 11-12

ring bonds :

1-8 1-9 8-11 9-10 10-11

exact/norm bonds :

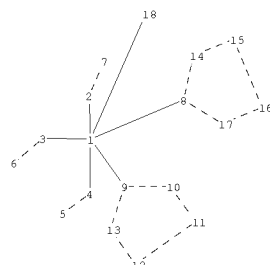
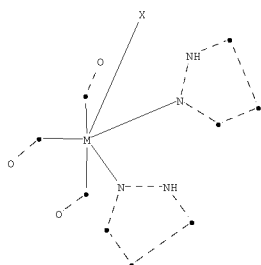
1-8 1-9 2-7 3-6 4-5 8-11 9-10 10-11 11-12

exact bonds :

1-2 1-3 1-4 10-13

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom
12:CLASS 13:CLASS



chain nodes :

1 2 3 4 5 6 7 18

ring nodes :

8 9 10 11 12 13 14 15 16 17

chain bonds :

1-2 1-3 1-4 1-8 1-9 1-18 2-7 3-6 4-5

ring bonds :

8-14 8-17 9-10 9-13 10-11 11-12 12-13 14-15 15-16 16-17

exact/norm bonds :

2-7 3-6 4-5 8-14 8-17 9-10 9-13 10-11 11-12 12-13 14-15 15-16 16-17

exact bonds :

1-2 1-3 1-4 1-8 1-9 1-18

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom
12:Atom 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:CLASS